GenCore version 5.1.6
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OM nucleic - nucleic search, using sw model

Run on: May 4, 2005, 12:02:42 ; Search time 292.117
Seconds

                                        (without alignments)
                                        1814.018 Million cell
updates/sec

Title: US-09-300-482-4
Perfect score: 263
Sequence: 1
gaagcacttttggatgttgc..........agnngnacntnnnnnganna 263

Scoring table: IDENTITY_NUC
                 Gapop 10.0 , Gapext 1.0

Searched: 9369276 seqs, 1007425964 residues

Total number of hits satisfying chosen parameters: 18738552

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
                 Maximum Match 100%
                 Listing first 45 summaries

Database :   Pending_Patents_NA_New:*
                 1: /cgn2_6/ptodata/1/pna/PCT_NEW_COMB.seq:*
                 2: /cgn2_6/ptodata/1/pna/PCT_NEW_COMB.seq2:*
                 3: /cgn2_6/ptodata/1/pna/US06_NEW_COMB.seq:*
                 4: /cgn2_6/ptodata/1/pna/US07_NEW_COMB.seq:*
                 5: /cgn2_6/ptodata/1/pna/US08_NEW_COMB.seq:*
                 6: /cgn2_6/ptodata/1/pna/US09_NEW_COMB.seq:*
                 7: /cgn2_6/ptodata/1/pna/US10_NEW_COMB.seq:*
                 8: /cgn2_6/ptodata/1/pna/US10_NEW_COMB.seq2:*
                 9: /cgn2_6/ptodata/1/pna/US10_NEW_COMB.seq3:*
                 10: /cgn2_6/ptodata/1/pna/US11_NEW_COMB.seq:*
                 11: /cgn2_6/ptodata/1/pna/US11_NEW_COMB.seq2:*
                 12: /cgn2_6/ptodata/1/pna/US11_NEW_COMB.seq3:*
                 13: /cgn2_6/ptodata/1/pna/US60_NEW_COMB.seq:*